## 100140/0483140≡ .IC13 Rec'd PCT/PT0 30 OCT 2001

## SEQUENCE LISTING <110> MIYATA, Toshio KUROKAWA, Kivoshi <120> Meg-3 protein <130> 2605/101 <140> Unknown <141> 2001-10-30 <160> 8 <170> FastSEQ for Windows Version 4.0 <210> 1 <211> 3768 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (53)..(2251) -400 - 1 caggaactgg gecageteeg gteeetteet tttggggete teactetgga gg atg ggg 58 Met Gly 1 tgg atg gga gaa aaa acc ggg aag atc ctg acg gag ttc ctc cag ttc 106 Trp Met Gly Glu Lys Thr Gly Lys Ile Leu Thr Glu Phe Leu Gln Phe tat gaa gac cag tat ggc gtg gct ctc ttc aac agc atg cgc cat gag 154 Tyr Glu Asp Gln Tyr Gly Val Ala Leu Phe Asn Ser Met Arg His Glu 25 att gag ggc acg ggg ctg ccg cag gcc cag ctg ctc tgg cgc aag gtg 202 Ile Glu Gly Thr Gly Leu Pro Gln Ala Gln Leu Leu Trp Arg Lys Val 35 45 cca ctg gac gag ege ate gte ttc teg ggg aac ete tte cag cac cag 250 Pro Leu Asp Glu Arg Ile Val Phe Ser Gly Asn Leu Phe Gln His Gln gag gac agc aag aag tgg aga aac cgc ttc agc ctc gtg ccc cac aac 298 Glu Asp Ser Lys Lys Trp Arg Asn Arg Phe Ser Leu Val Pro His Asn tac ggg ctg gtg ctc tac gaa aac aaa gcg gcc tat gag cgg cag gtc 346 Tyr Gly Leu Val Leu Tyr Glu Asn Lys Ala Ala Tyr Glu Arg Gln Val cca cca cga gcc gtc atc aac agt gca ggc tac aaa atc ctc acg tcc 394

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110

Pro Pro Arg Ala Val Ile Asn Ser Ala Gly Tyr Lys Ile Leu Thr Ser

105

100

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<sup>&</sup>lt;210> 2 <211> 733

<sup>&</sup>lt;212> PRT <213> Homo sapiens

<sup>-400- 2</sup> 

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Gln Phe Tyr Glu Asp Gln Tyr Gly Val Ala Leu Phe Asn Ser Met Arg
20 25 30

His Glu Ile Glu Gly Thr Gly Leu Pro Gln Ala Gln Leu Leu Trp Arg  $35 \hspace{1cm} 40 \hspace{1cm} 45$ 

Lys Val Pro Leu Asp Glu Arg Ile Val Phe Ser Gly Asn Leu Phe Gln 50 60

His Gln Glu Asp Ser Lys Lys Trp Arg Asn Arg Phe Ser Leu Val Pro

His Asn Tyr Gly Leu Val Leu Tyr Glu Asn Lys Ala Ala Tyr Glu Arg Gln Val Pro Pro Arg Ala Val Ile Asn Ser Ala Gly Tyr Lys Ile Leu 105 Thr Ser Val Asp Gln Tyr Leu Glu Leu Ile Gly Asn Ser Leu Pro Gly Thr Thr Ala Lys Ser Gly Ser Ala Pro Ile Leu Lys Cys Pro Thr Gln 135 Phe Pro Leu Ile Leu Trp His Pro Tyr Ala Arg His Tyr Tyr Phe Cys Met Met Thr Glu Ala Glu Gln Asp Lys Trp Gln Ala Val Leu Gln Asp Cys Ile Arg His Cys Asn Asn Gly Ile Pro Glu Asp Ser Lys Val Glu Gly Pro Ala Phe Thr Asp Ala Ile Arg Met Tyr Arg Gln Ser Lys Glu 200 Leu Tyr Gly Thr Trp Glu Met Leu Cys Gly Asn Glu Val Gln Ile Leu Ser Asn Leu Val Met Glu Glu Leu Gly Pro Glu Leu Lys Ala Glu Leu 235 Gly Pro Arg Leu Lys Gly Lys Pro Gln Glu Arg Gln Arg Gln Trp Ile Gln Ile Ser Asp Ala Val Tyr His Met Val Tyr Glu Gln Ala Lys Ala Arg Phe Glu Glu Val Leu Ser Lys Val Gln Gln Val Gln Pro Ala Met 280 Gln Ala Val Ile Arg Thr Asp Met Asp Gln Ile Ile Thr Ser Lys Glu Leu Leu Ala Ser Lys Ile Arg Ala Phe Ile Leu Pro Lys Ala Glu Val Cys Val Arg Asn His Val Gln Pro Tyr Ile Pro Ser Ile Leu Glu Ala Leu Met Val Pro Thr Ser Gln Gly Phe Thr Glu Val Arg Asp Val Phe 345 Phe Lys Glu Val Thr Asp Met Asn Leu Asn Val Ile Asn Glu Gly Gly 355 360 365

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380

Ile Asp Lys Leu Gly Glu Tyr Met Glu Lys Leu Ser Arg Leu Ala Tyr

375

370

His Pro Leu Lys Met Gln Ser Cys Tyr Glu Lys Met Glu Ser Leu Arg 385 Leu Asp Gly Leu Gln Gln Arg Phe Asp Val Ser Ser Thr Ser Val Phe Lys Gln Arg Ala Gln Ile His Met Arg Glu Gln Met Asp Asn Ala Val Tyr Thr Phe Glu Thr Leu Leu His Gln Glu Leu Gly Lys Gly Pro Thr 440 Lys Glu Glu Leu Cys Lys Ser Ile Gln Arg Val Leu Glu Arg Val Leu Lys Lys Tyr Asp Tyr Asp Ser Ser Ser Val Arg Lys Arg Phe Phe Arg 475 Glu Ala Leu Leu Gln Ile Ser Ile Pro Phe Leu Leu Lys Lys Leu Ala Pro Thr Cvs Lvs Ser Glu Leu Pro Arg Phe Gln Glu Leu Ile Phe Glu Asp Phe Ala Arg Phe Ile Leu Val Glu Asn Thr Tyr Glu Glu Val Val Leu Gln Thr Val Met Lys Asp Ile Leu Gln Ala Val Lys Glu Ala Ala Val Gln Arg Lys His Asn Leu Tyr Arg Asp Ser Met Val Met His Asn Ser Asp Pro Asn Leu His Leu Leu Ala Glu Gly Ala Pro Ile Asp Trp Gly Glu Glu Tyr Ser Asn Ser Gly Gly Gly Gly Ser Pro Ser Pro Ser Thr Pro Glu Ser Ala Thr Leu Ser Glu Lys Arg Arg Arg Ala Lys Gln Val Val Ser Val Val Gln Asp Glu Glu Val Gly Leu Pro Phe Glu Ala Ser Pro Glu Ser Pro Pro Pro Ala Ser Pro Asp Gly Val Thr Glu Ile Arg Gly Leu Leu Ala Gln Gly Leu Arg Pro Glu Ser Pro Pro Pro Ala Gly Pro Leu Leu Asn Gly Ala Pro Ala Gly Glu Ser Pro Gln Pro Lys Ala Ala Pro Glu Ala Ser Ser Pro Pro Ala Ser Pro Leu Gln His Leu 675 680 685

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Leu Pro Gly Lys Ala Val Asp Leu Gly Pro Pro Lys Pro Ser Asp Gln
                        695
                                             700
Glu Thr Gly Glu Gln Val Ser Ser Pro Ser Ser His Pro Ala Leu His
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□ Page(s)of	were not
present for scanning.	(Document title)

& Scanned copy is best available. Figure 2 is Very Dark